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Fig. 1

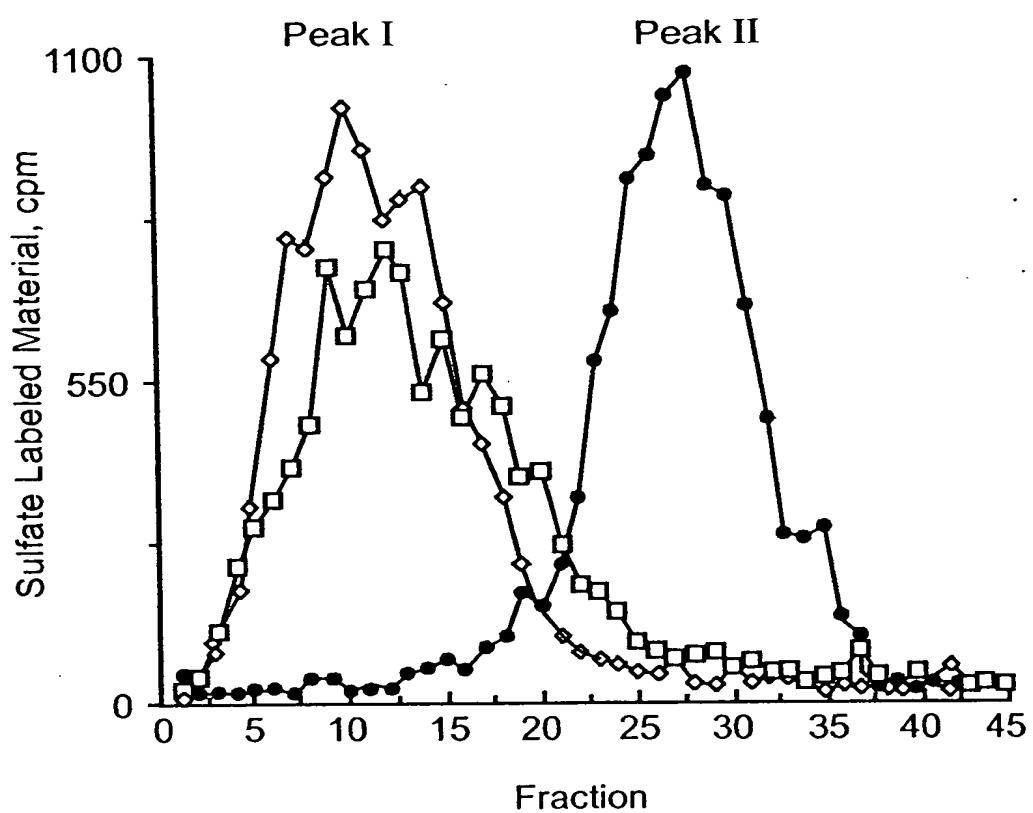
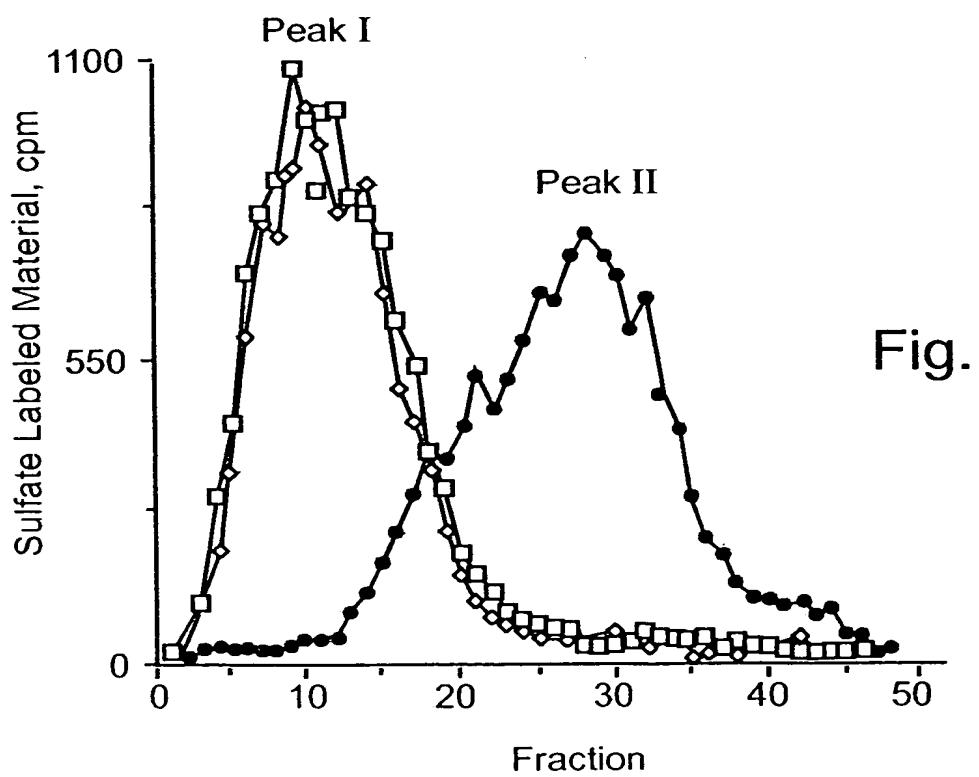
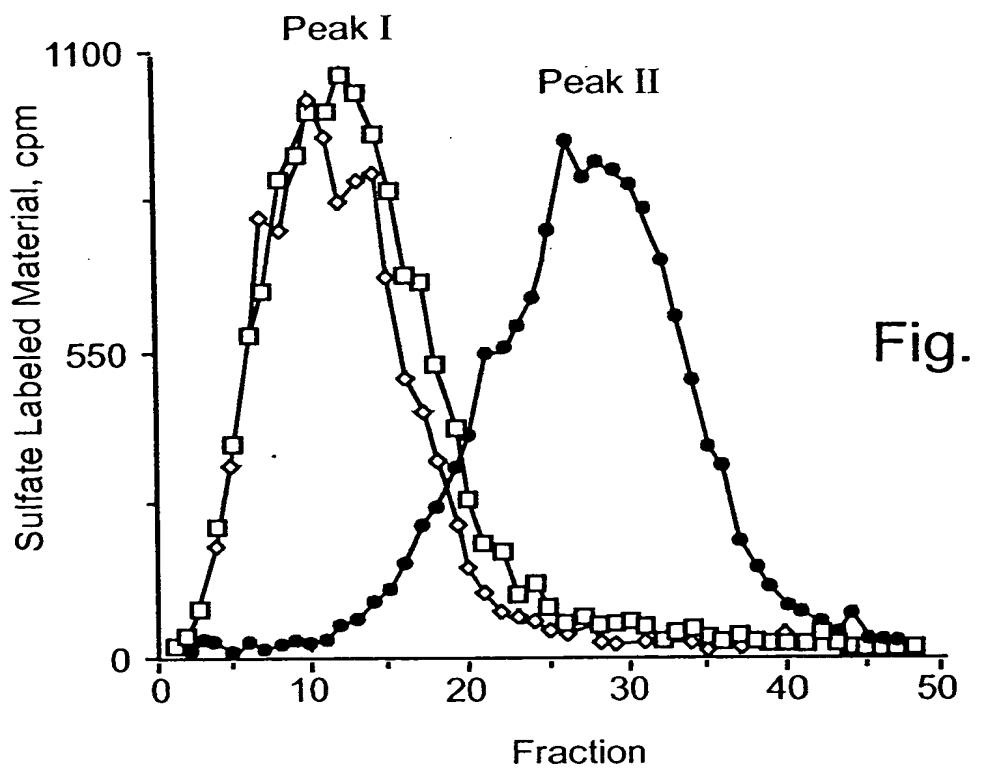


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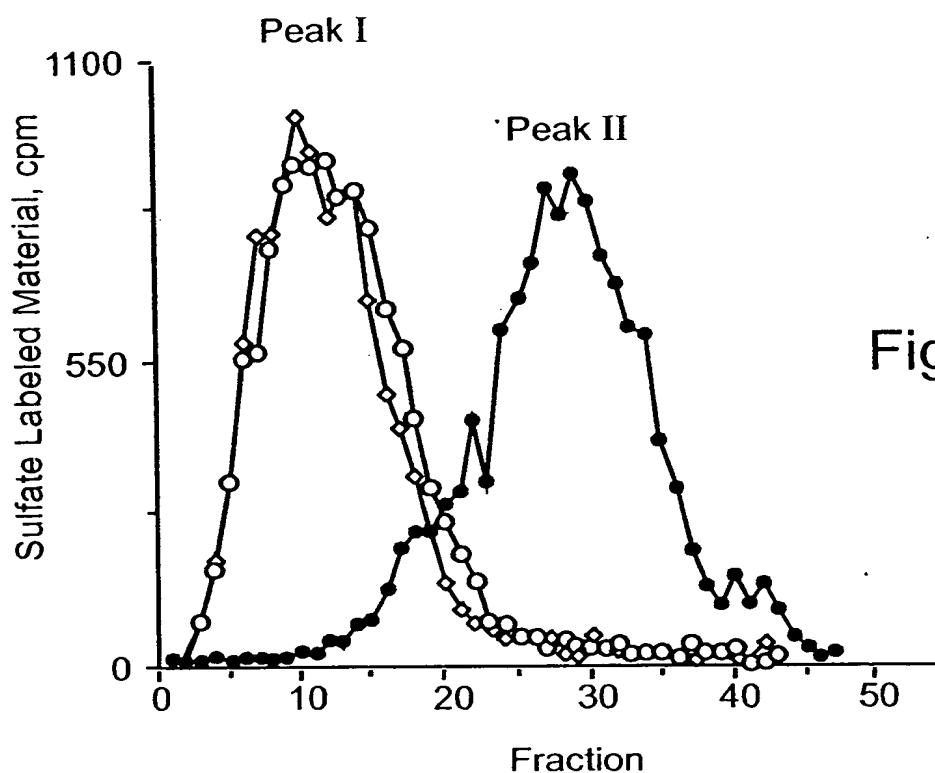


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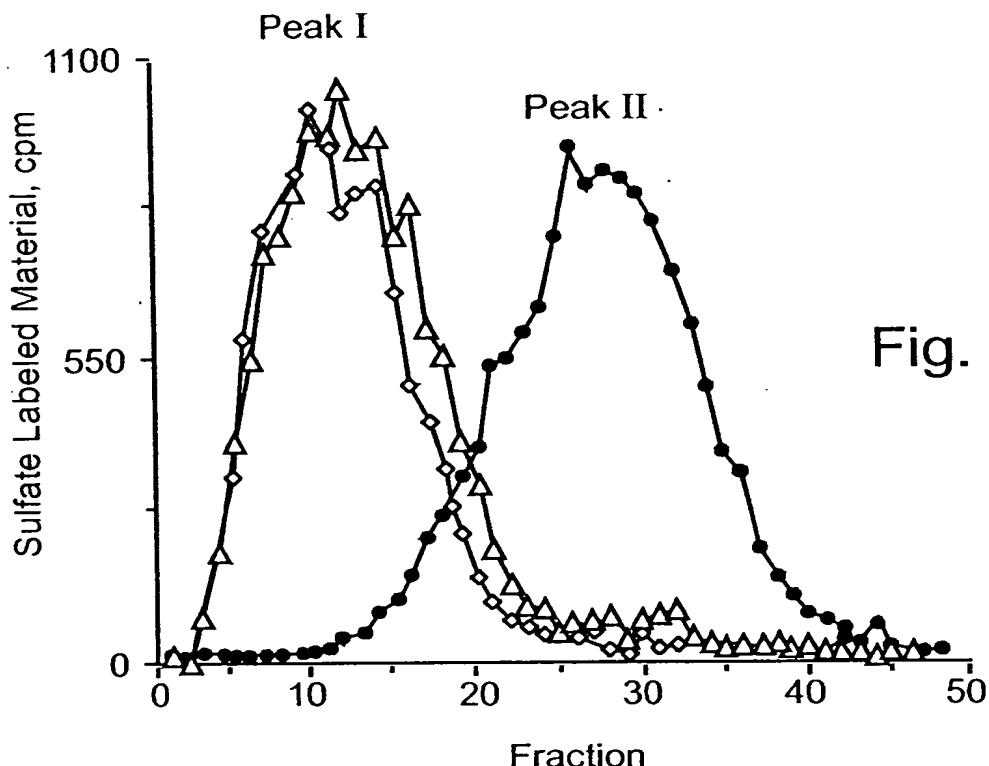


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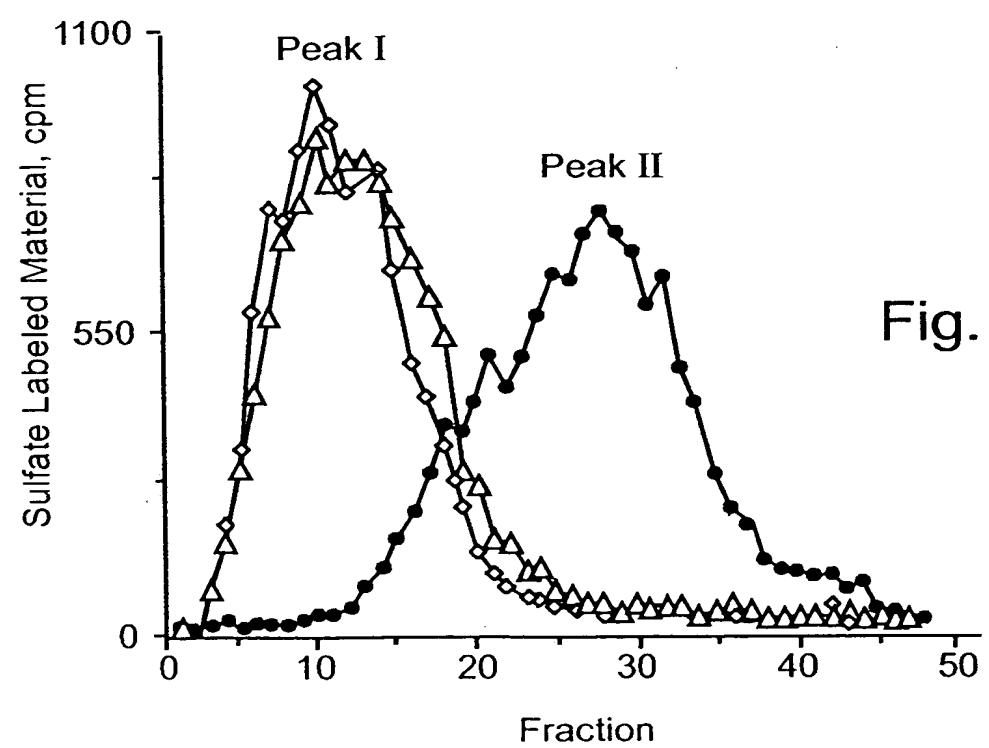
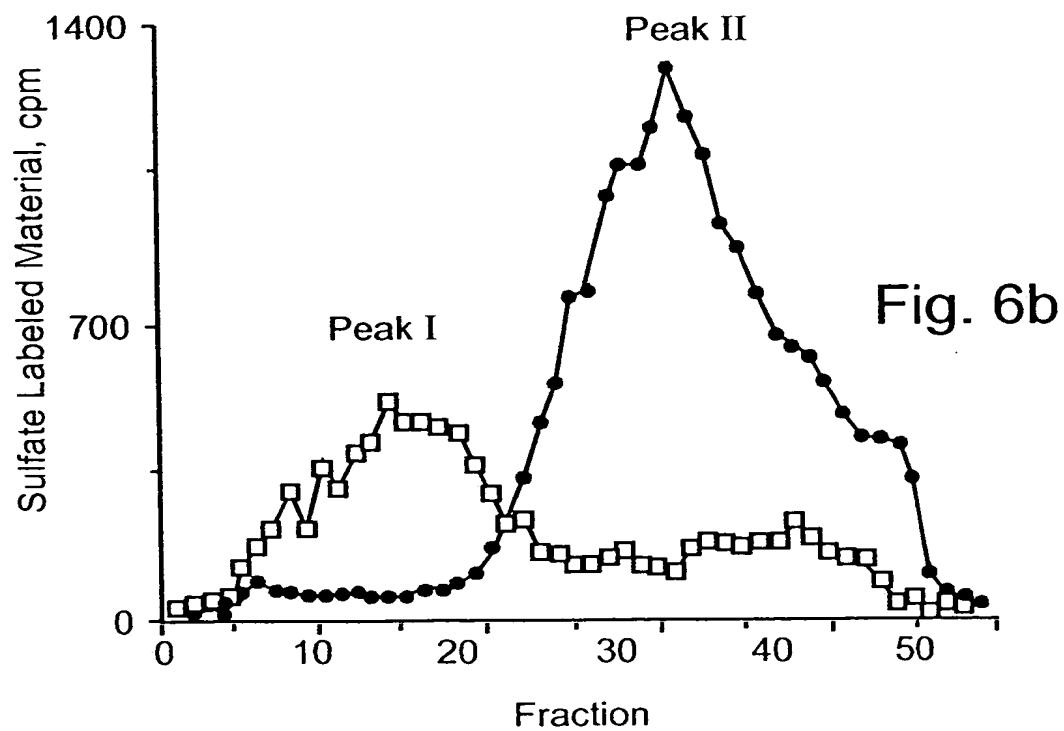
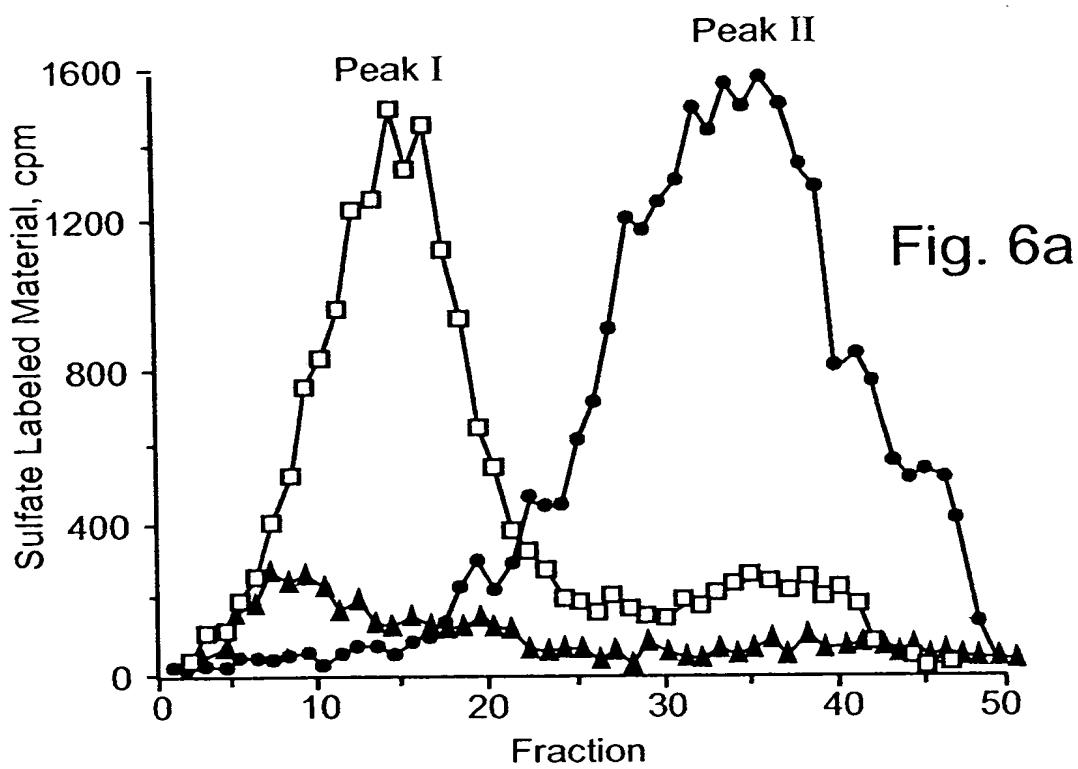
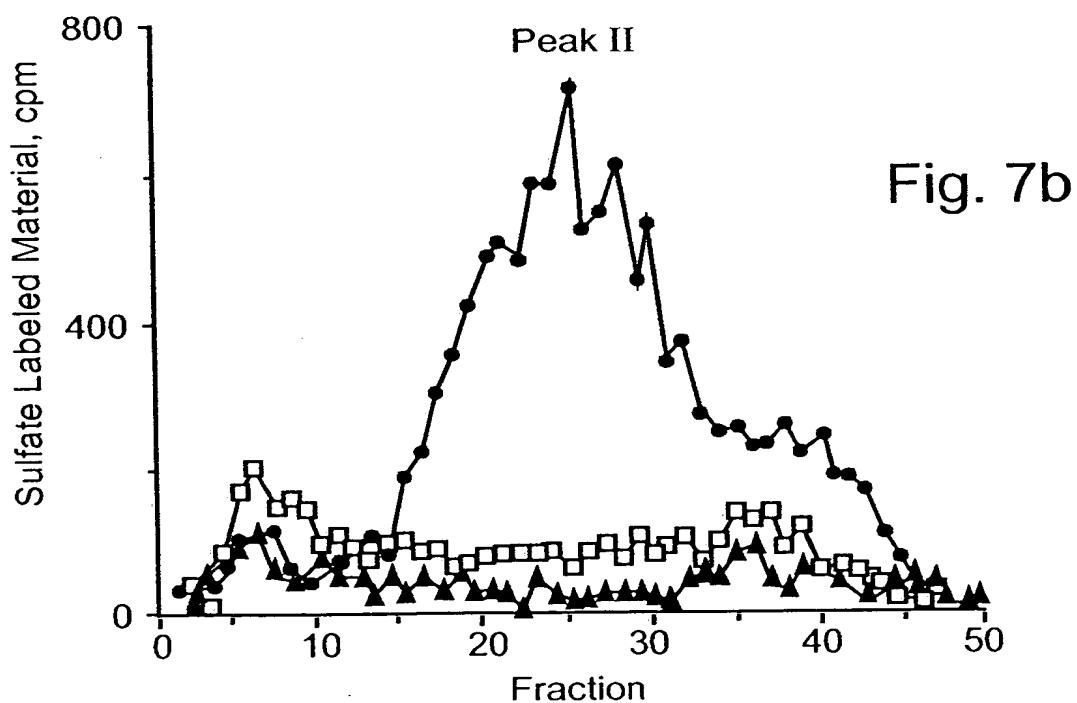
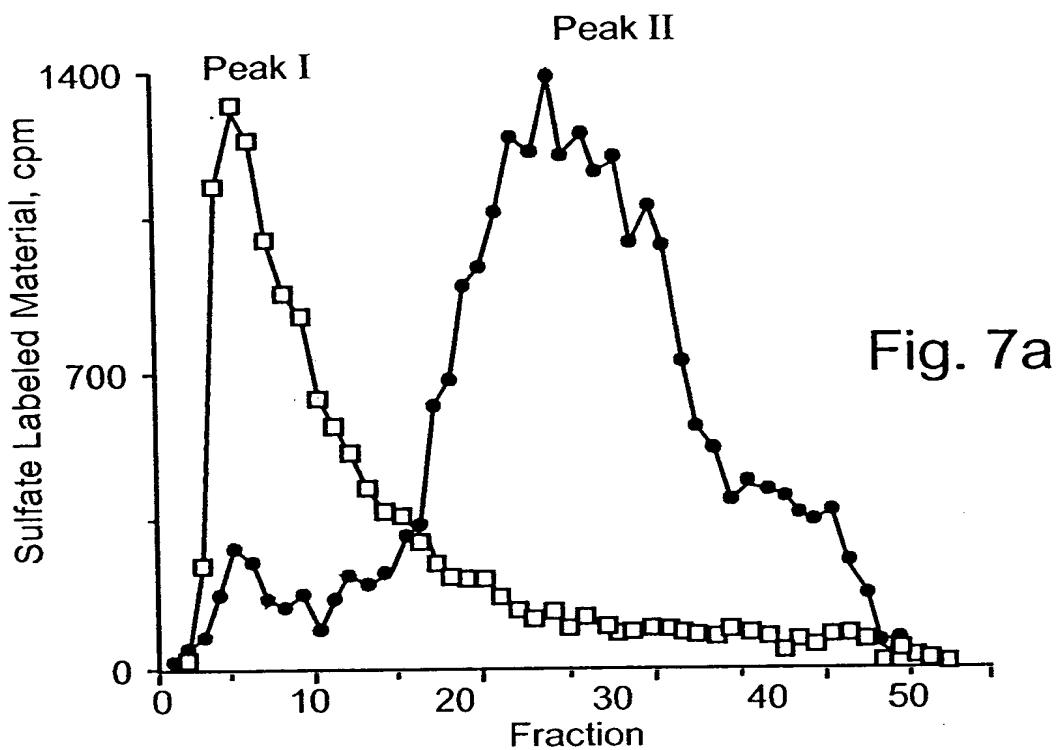
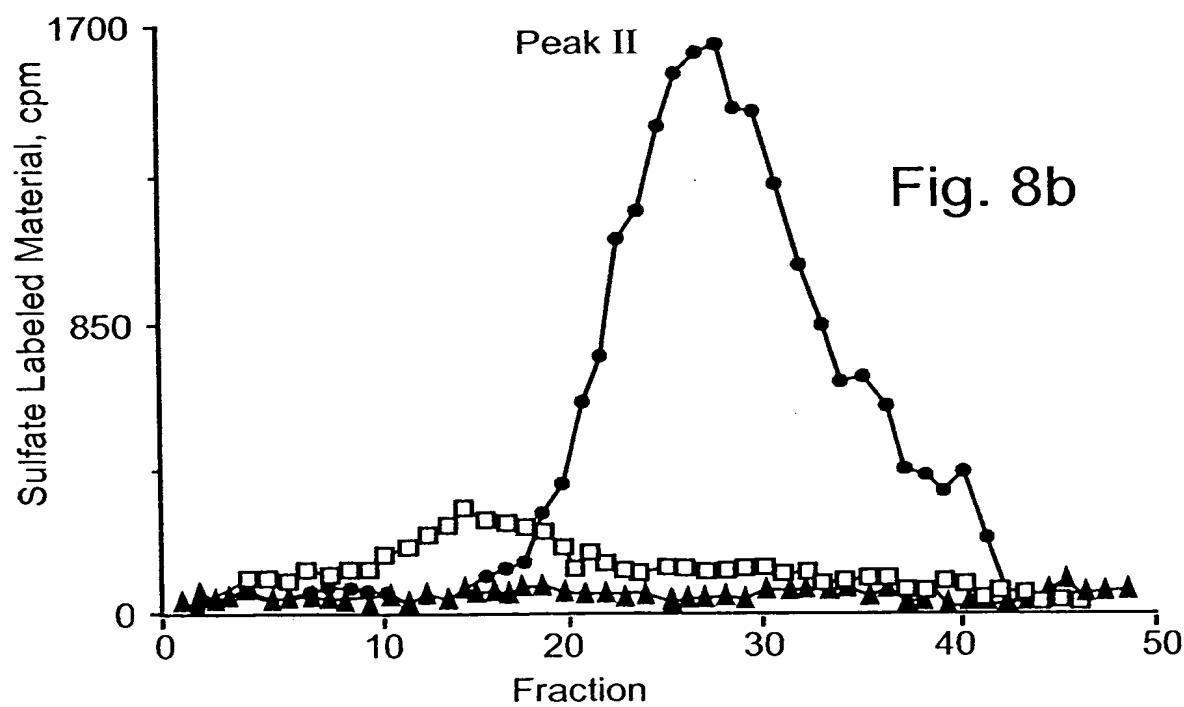
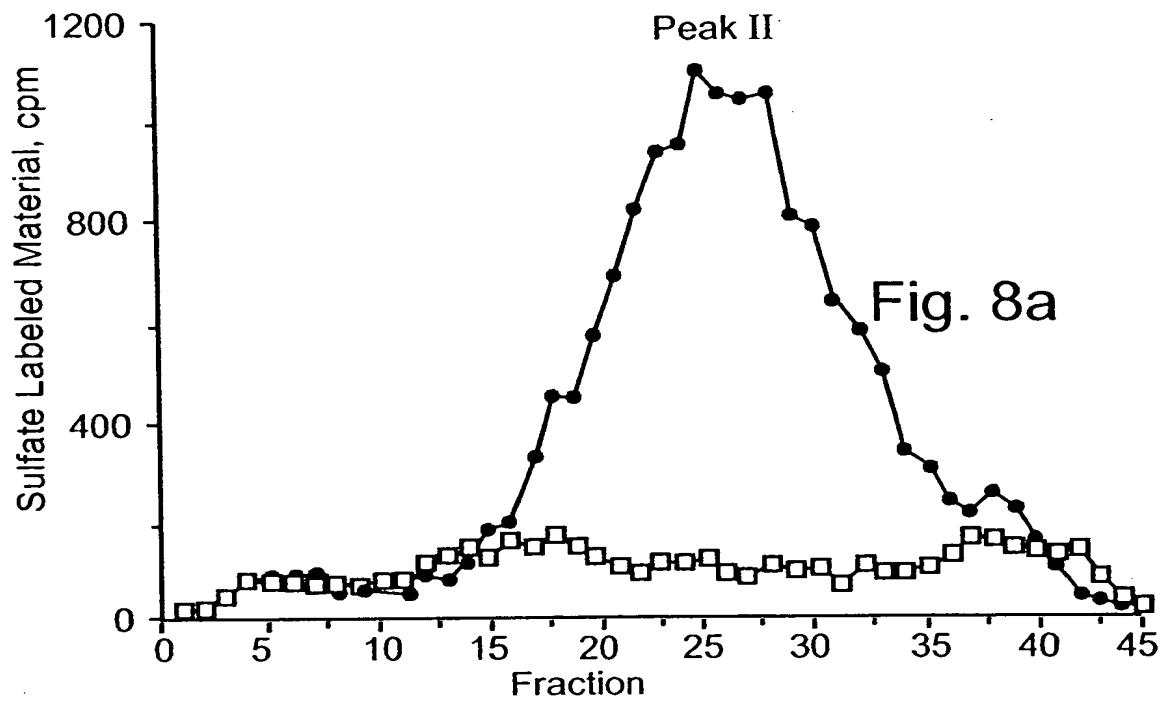
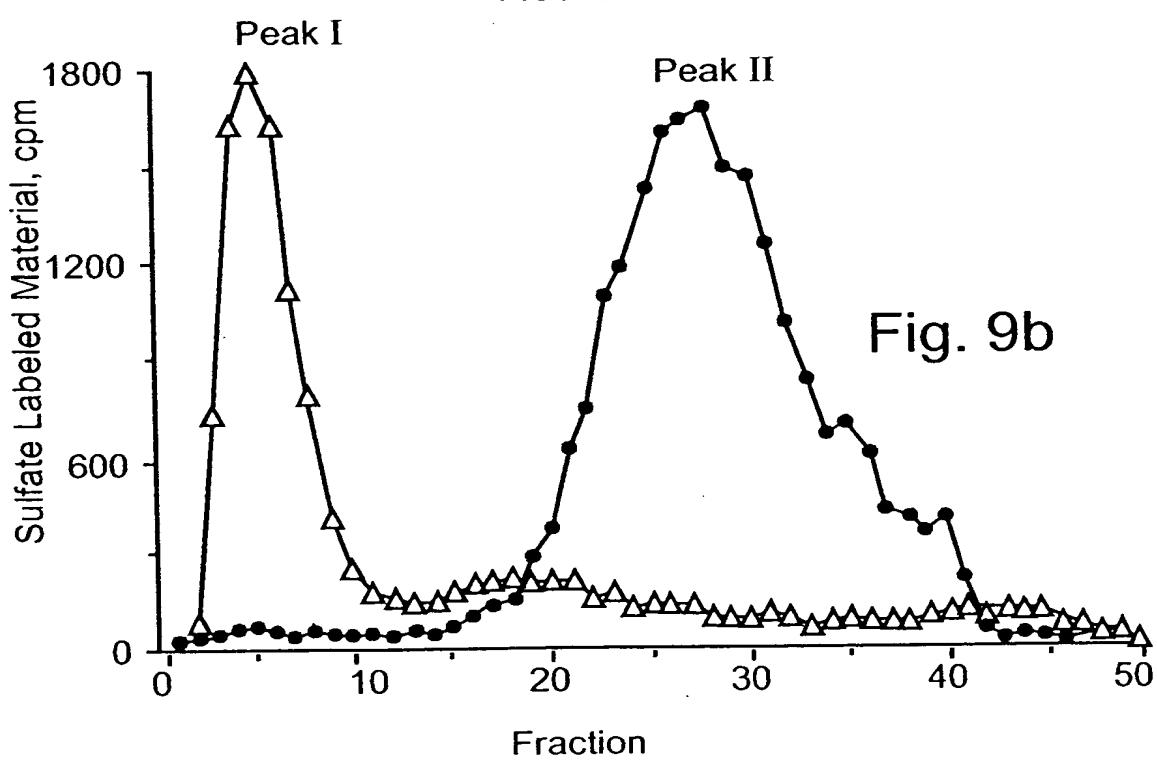
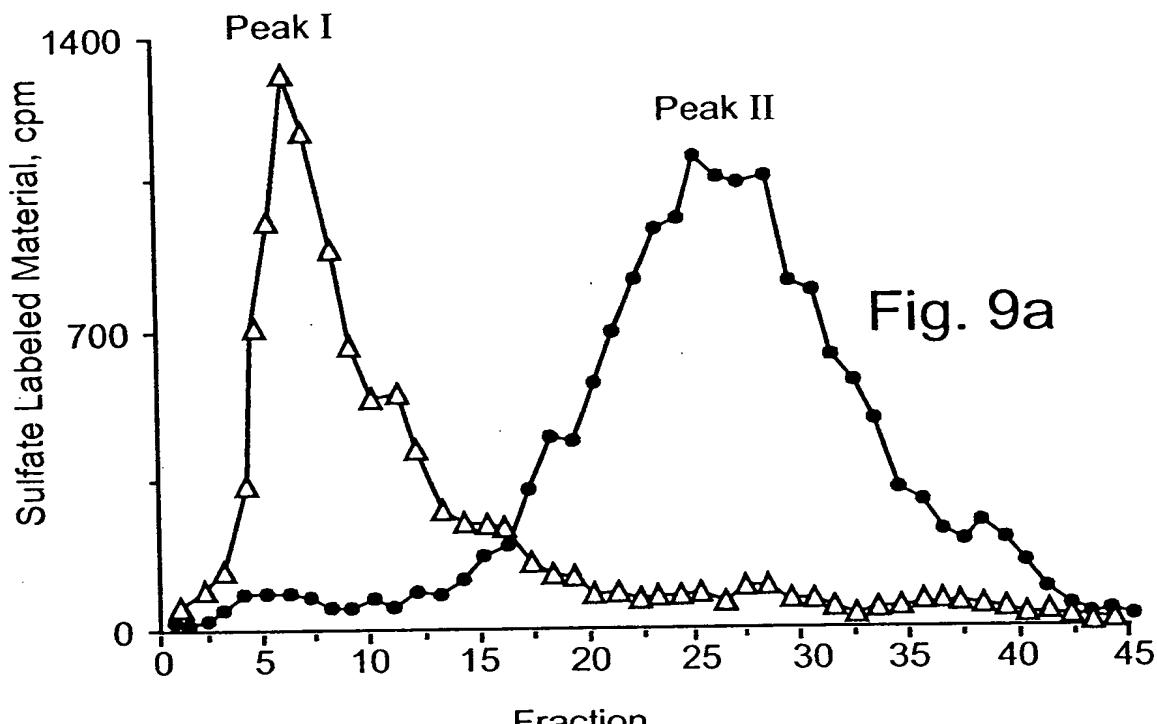


Fig. 5b









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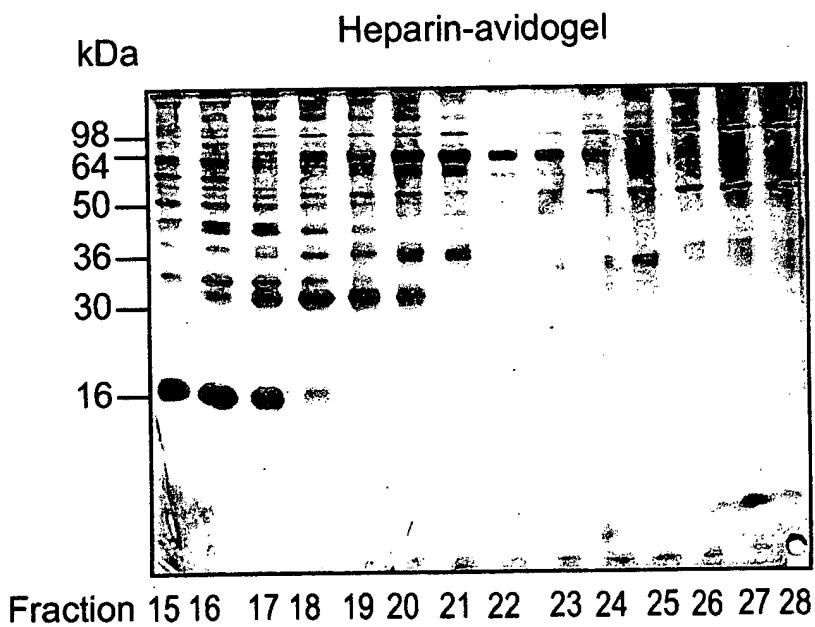
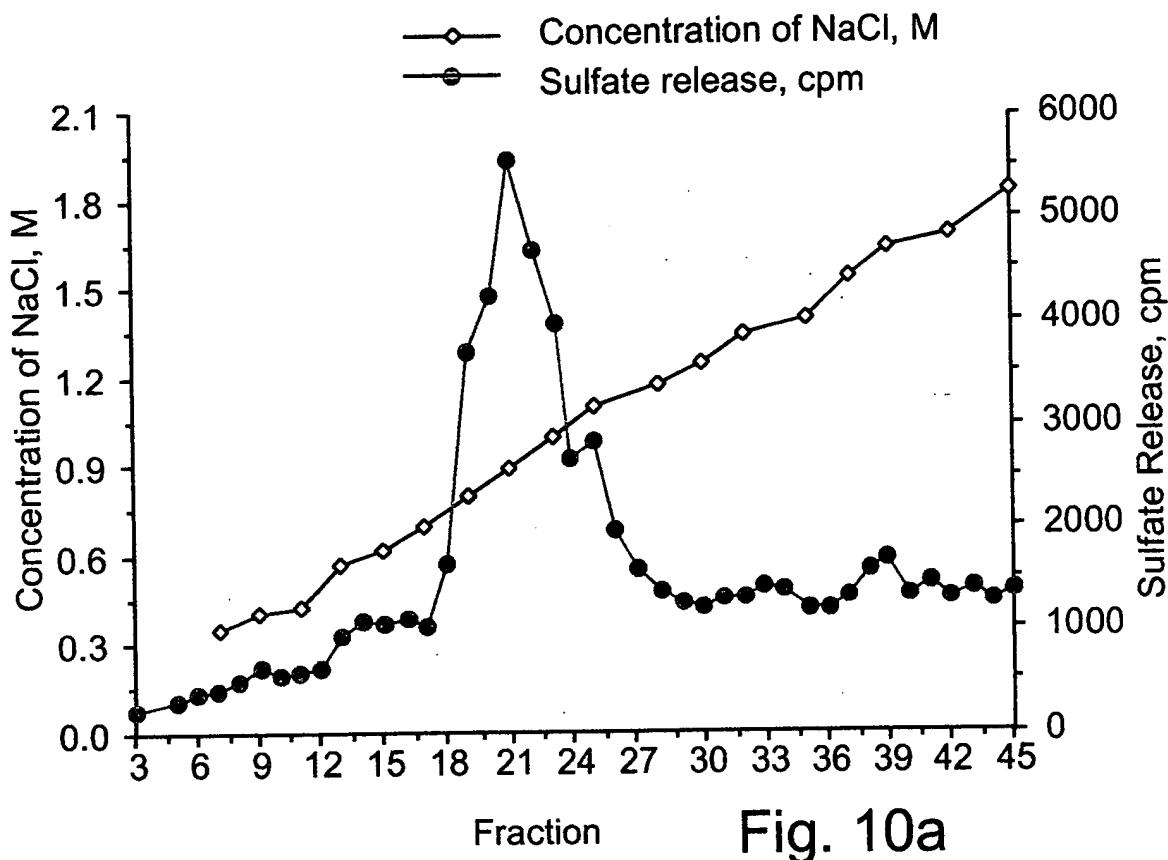


Fig. 10b

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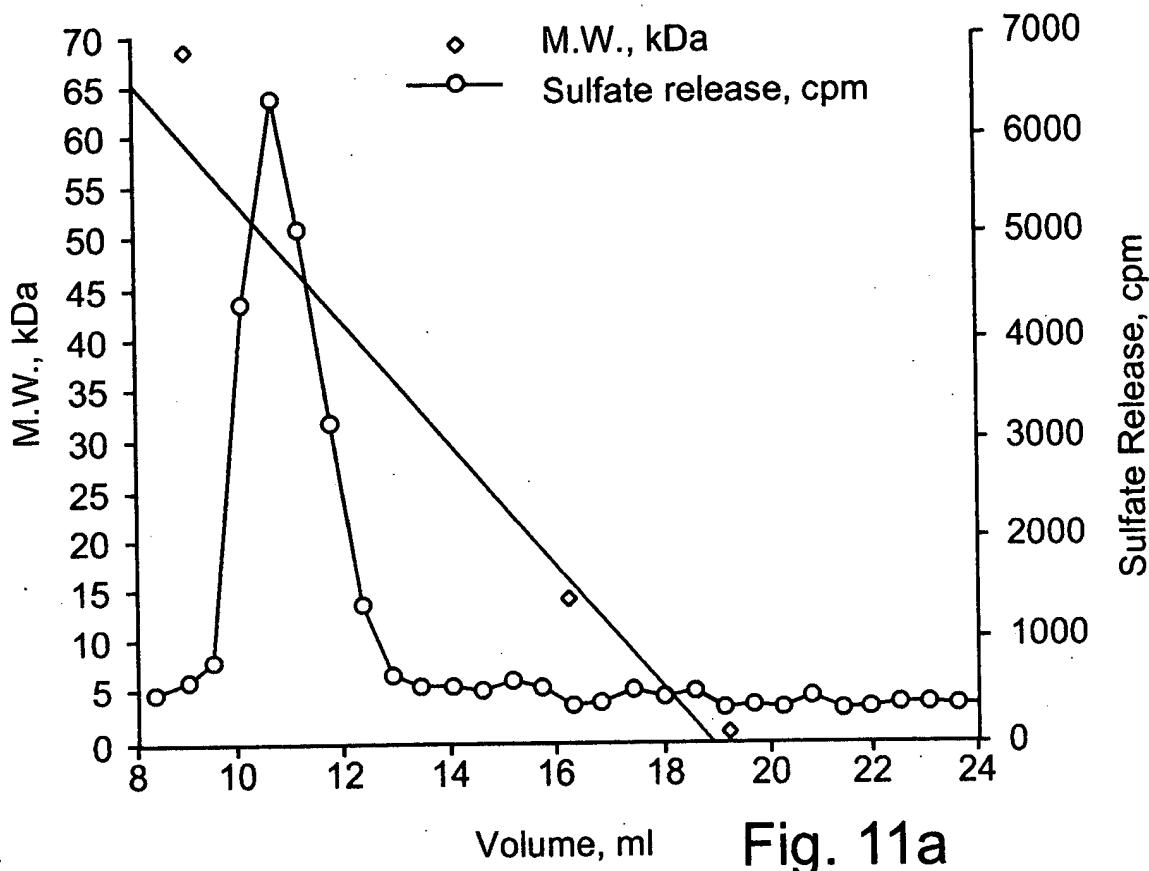


Fig. 11a

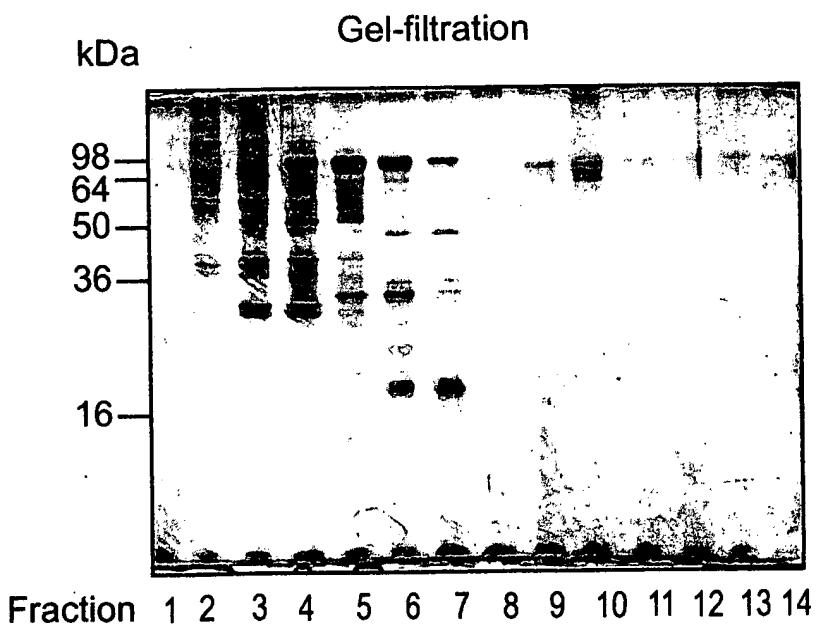


Fig. 11b

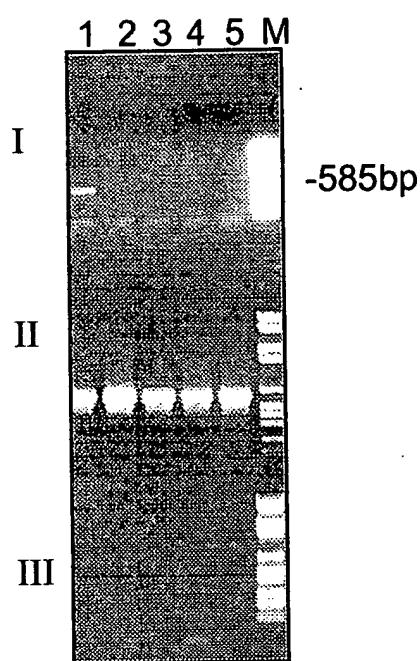


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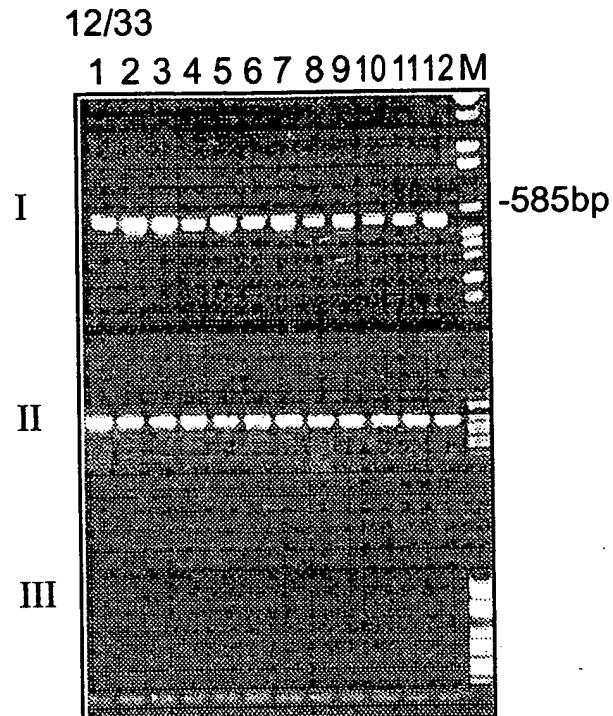


Fig. 12b

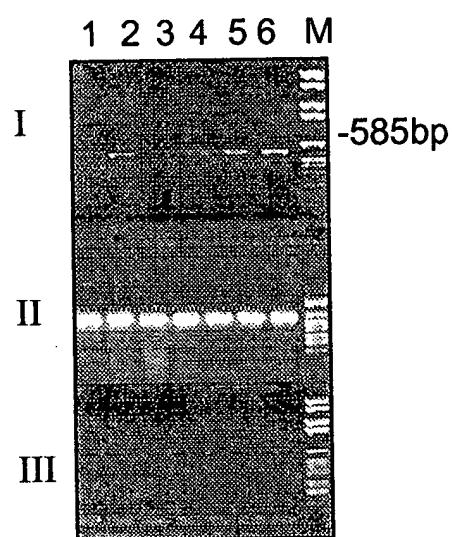


Fig. 12c

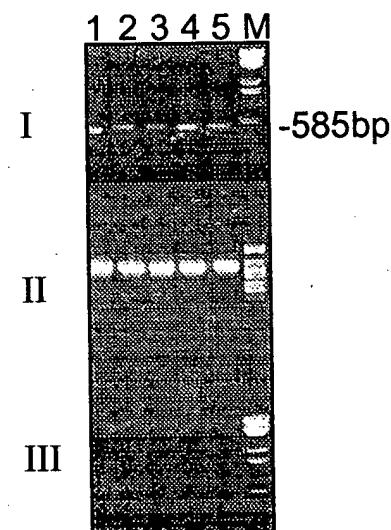


Fig. 12d

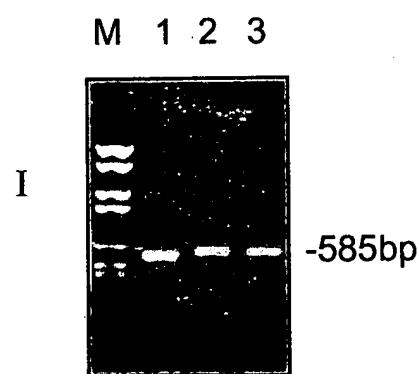


Fig. 12e

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Fig. 13

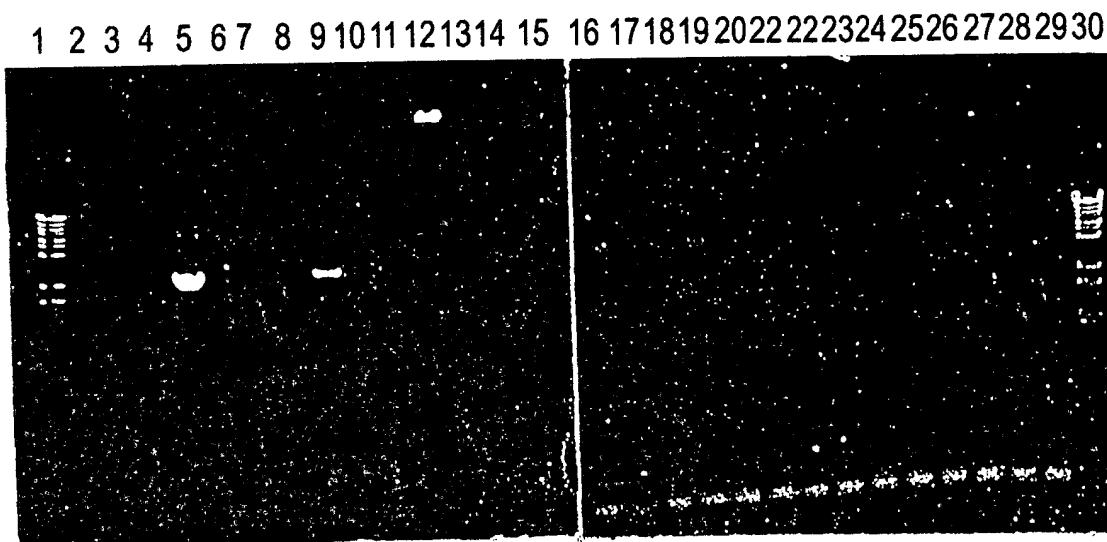


Fig. 14

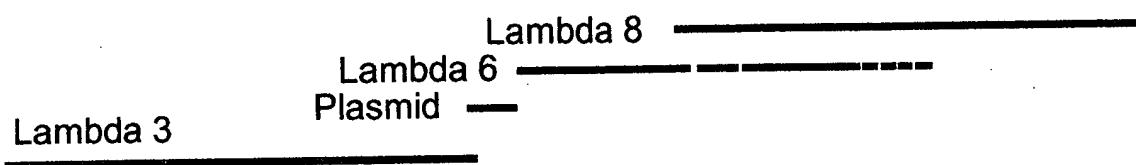
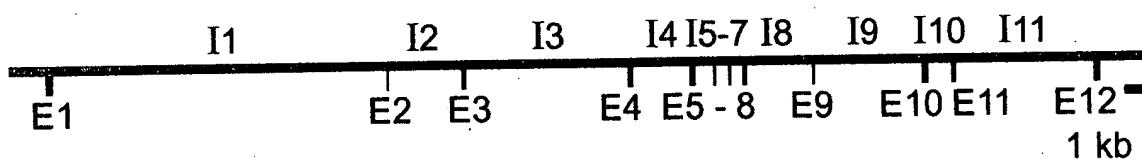


Fig. 15

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Fig. 16

Fig. 16
(continued)

Fig. 16
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aactagaatttcttcagttacaacacagaatagtattcattatgaaa	24150
gcgaacttggaggccttcatgtggtgcacatctaaccattaaatgtga	24200
cgttttttttttagGAAGCTCTGTAGATGTGCTATACACTTTGCACAACT	24250
R S S V D V L Y T F A N	
GCTCAGGACTGGACTTGTATCTTGGCTAAATGCCTTATTAAGAACAGCA	24300
C S G L D L I F G L N A L L R T A	24350
GATTTGCAGTGGAACAGTCTTAATGCTCAGTTGCTCCTGGACTACTGCTC	
D L Q W N S S N A Q L L L D Y C S	
TTCCAAGGGGTATAACATTCTGGAACTAGGCAATGgtgagtacccca	24400
S K G Y N I S W E L G N	
gggaaacaaattcataataaaggagattccccactagcattattttttttt	24450
tttcttt	24500
tgcccaaggctggagtgcagtggcgccacctcgctacttgaagctctgc	24550
ctcccaaaacgcatttcctgcctcagcctcccgagtagtgggactac	24600
aggcaccgcaccccgccccgttaatttttttttttttttttttttttttt	24650
tttttttgcatttttagtagagacggggttcacctgttagccaggatg	24700
gtcttgatctcctgacccctgtatctgccttctcggcctccaaagtgc	24750
tgggattacaggcgtgagccaccaggccccgttagcattattttttatga	24800
cacttt	24850
agtgcagtggcgccatctcggtcaactgcaagctccacccctccagggtca	24900
cgccatttcctgcctcagcctcccgagtagtggactacacgcaccccg	24950
ccaccacgcccggtaattttttgtattttttagtagagacggggttca	25000
ccgtgttagccagatggctctatccatgtaccatgtatctgcccggcc	25050
tcggccctccaaagtgggattacaggcgtgagccactgcggccggcc	25100
aacacttt	25150
caagtgtcaacaatgcacattttggaaagtgcattggcagaaacttcctg	25200
ctgtatattttccagaaaccttattttgttaatcccaagtttatgttacatt	25250
tgaagtgagaaccagggtggccagcaacgttcccaagctccaaagtccc	25300
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gtttctcaatgtgtcagttaatctcagtaaccatgtcaacccatttca	25450
acctgcccacccgttcaagaacttgcacgtatagaatcttaactgtgggtca	25500
agctcctgactgtctcccttcacttttttttttttttttttttttttttt	25550
ttttactataagtattcatgtattgcacatttttttttttttttttttttt	25600
gctttttccacatatcagccaatggaaaataaggattaaatggaaatgaa	25650
atgttagtaataggataagcacaagtcttcctgtcttttttttttttttt	25700
tt	25750
cagacaagattt	25800
ggcgtgttcatagctcaatgttaccccttcaactcctgggctatgcacat	25850
ctcacacccctcagccccctgattagcttaggactacactatgcctagccat	25900
tt	25950
ctcaagtaatccctgcctcgccctctaaagtgtggattatagcata	26000
tgagccactgtggccggctcaaaacccctttttttccaaagtaatggatt	26050
attagatatggaaatatactgttgcctccagatatccatcatccatgttt	26100
attaccctcattatataacttcaaaatttttttttttttttttttttttttt	26150
ttatacagttaaaatt	26200
ctatgagtt	26250
ctctgtcactcaggctggagtgcgggtgcgtatggctactatggc	26300
ctcgacccctctggctcaagtgcattccctccctcagccctccaaactgttag	26350
actacaggcatgcaccacacatctagctaatttttttttttttttttttttt	26400
acaaggctttactatgttacccagagtggctcaaaactctggcctca	26450
qqqatcttcgtctcagccatccaaatgttgcggattacaggcatgagc	

Fig. 16
(continued)

catagcgccagacacctggttactttcttgacttgaattacaagttt	26500
tgttaattggaaaatgtttgtgtttaaatactgctgtatgtttgtct	26550
tttaaatacaacattctcgatataatatttgagaattgctgtttcaag	26600
AACCTAACAGTTCTTAAGAAGGCTGATATTTCATCAATGGGTGCGAG	26650
E P N S F L K K A D I F I N G S Q	
TTAGGAGAAGATTATTCAATTGCATAAACTCTAAGAAAGTCCACCTT	26700
L G E D F I Q L H K L L R K S T F	
CAAAAATGCAAAACTCTATGGCCTGATGTTGGTCAGCCTCGAAGAAAGA	26750
K N A K L Y G P D V G Q P R R K	
CGGCTAAGATGCTGAAGAGgttaggaactagaggatgcagaatactttac	26800
T A K M L K S	
ttttcttcttttctttgagacagagtctcaactctgtcagccagactg	26850
gagtgcagtggtaacaatcatggctactgcaacttcgacccaggctc	26900
aagcaatctccatctcagccccacaatagctggactacagggtgcac	26950
atcaccacacccctggctactttaaaaaaattttttagagatgggtct	27000
ccctgtgttgccaggctggctcttgaattcctgtgtcaagccatccc	27050
cccacccagccccccagggtccaggattacaggcatgagccaccacac	27100
ccagccaccactttcttaaaaaaaaaaaaagattctctgttagaccaa	27150
tcctcaatagtccacatgttattaaacaatctgctgcctgaatacatgt	27200
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gcaaatgtcacatgtataaaaaatttgcataaaaaatttaggaattttgt	27300
ttacctgatctaaagcagtaaccagccatttcttagggataaaaactct	27350
catgcgtatattgtgcataatattgtactgtactgtgataataaaa	27400
atttttttctagCTTCCTGAAGGCTGGTGGAGAAGTGATTGATTCAAGTT	27450
F L K A G G E V I D S V	
ACATGGCATCAgtaagtatgtctctttaataacttagggaaagtaagg	27500
T W H H	
ctagcttttatttattaccttagtattcaaaaagttagttcattaactg	27550
aattgactgcagttcaataagaaacaaaatagtgtctcaagtgcactgt	27600
actccaaattttatattaaataaaaaattttaaatataatgt	27650
tagtggttctataaaagatcaacttatacagaagaacacagtgcaccaattaa	27700
ccatggAACatataaagttagctaaaaccaattgtctgcaccaaggat	27750
acccaggagtagatgtccttgcactgtgttttcaagacagagtaact	27800
gattttctagttacttgcataatggactctcctcataactcccttcca	27850
tcttggctttccctagtagaaacttctaccttttttagtaacaggtag	27900
tgggagaggtaaagaaggagaataaggtcagcaattaacctaaaagcagaa	27950
agtaaaaattttgttatttttctgaatattttctgtgtatatttagCTAC	28000
Y	
TATTTGAATGGACGGACTGCTACCAGGGAAAGATTCTAAACCCCTGATGT	28050
Y L N G R T A T R E D F I N P D V	
ATTGGACATTTTATTCATCTGTGCAAAAAGTTCTCAGgtaatagtct	28100
L D I F I S S V Q K V F Q	
ttttaaacttttaatgtaaaaccagaatccttattttatagtctagta	28150
gttctaaattctataggtagtatattcatgtttttctaaatttttagag	28200
aacaaggactatgacttattccactgttagttttcccttagcattgggtc	28250
ttaccccatgtacgtgatttagaaatttggaaatattccaaatagcctttag	28300
tagaattaactcacatagatgataagaatgggtgggtcacttcatgttc	28350
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aagtttgaatgtgttaatcttcaacaccacagttgaaaccacaggta	28500
gttttttgcattaccatggatactttctgttctatagGTGGTTGAGA	28550
V V E	
GCACCAGGCCCTGGCAAGAAGGTCTGGTTAGGAGAAACAAAGCTCTGCATAT	28600
S T R P G K K V W L G E T S S A Y	
GGAGGGCGGAGCGCCCTTGCATCCGACACCTTGCAGCTGGCTTTATgtg	28650
G G G A P L L S D T F A A G F M	
agtgaagcagcgctggccttagggtaggtcagagtgcagcttctccatct	28700
tctattctgtgaaatagctcccaagccaaaaagcagatcaaagaccgtt	28750
tcagttggctgagccccaaaattcatgccagatttgcagaaaaatgattt	28800
actaaagcttgaggacatcttaacaagtttgcagatcaattactata	28850
aggatgtatgtttcagaaattttggccctttaattatggccctaaat	28900

Fig. 16
(continued)

ctttgcagcaataatatgtgagaggacagattgttagatatgtatgtat	43450
aaaaaatggtaatgacaattcagaggcgaggagattctgtaaacttaaa	43500
attactataatgaaattgatttgcagaggataatgttagaaaacac	43550
ccaatacctataactgtctgttaatgcttgccttctaccttctt	43600
ccttgtttcagttggaaagctttggctgcaagtaacagaaactcctaata	43650
tcaaatggcttaagcaataaggaaatgtatattccacataactagacgt	43700
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tctggtagcatgtggctgttagctgtttcatggcccttcaaaccctcat	43850
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tgaataactcttttcagagcttcacagcaaacccttcacgtctc	43950
ctcatgtcttattgttcagaaatggtaatgtggccatttcaccagtac	44000
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tggagagggtgttggtcagtctacaaaactgaacactgcagttctgcgtt	44100
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ttcaaatgtatgcgttatggatatagtatcttaaaatttttatttt	44200
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aaaattaaatttagatcttcaataactcttaattttatatgttaagtgg	44700
ttttatatttcacattgaaataaagtaattttataacccttgatatt	44750
gtatgactatttttagtaatgtaaagccctacagactcctacatttqga	44800
accactagtgtttcacccttgcattactatcaggatcctcga	44898

Fig. 16
(continued)

human	MLLRSKPALP	PPLMLLLGP	LGPLSPGALP	RPAQAQDVVD	LDFFTQEPLH	
mouse	~~~~~	ML	RLLLLWLWGP	LGALAQQAPA	GTAPTDVVD	LEFYTKRPLR
rat	~~~~~	~~~~~	~LLLLWLWGR	LRALTQGTPA	GTAPTDVVD	LEFYTKRQLFQ
						100
human	LVSPSFSLVT	IDANLATDPR	FLILLGSPKL	RTLARGLSPA	YLRFGGTKTD	
mouse	SVSPSFSLSI	IDASLATDPR	FLTFLGSPRL	RALARGLSPA	YLRFGGTKTD	
rat	SVSPSFSLSI	IDASLATDPR	FLTFLSSPRL	RALSRLGLSPA	YLRFGGTKTD	
						150
human	FLIFDPKES	TFEERSYWQS	QVNQDICKYQ	SIPPDVEEKL	RLEWPYQEQL	
mouse	FLIFDPDKEP	TSEERSYWKS	QVNHDICRSE	PVSAAVLRKL	QVEWPFQELL	
rat	FLIFDPNNEP	TSEERSYWQS	QDNNDICGSD	RVSADVL	~~~~~	
						200
human	LLREHYQKKEF	KNSTYSRSSV	DVLYTFANCS	GLDLIFGLNA	LLRTADLQWN	
mouse	LLREQYQKEF	KNSTYSRSSV	DMLYSFAKCS	GLDLIFGLNA	LLRTPDLRWN	
rat	~~~~~	~~~~~	~~~~~	~~~~~	~~~~~	
						250
human	SSNAQLL LDY	CSSKGYNISW	ELGNEPNSFL	KKADIFINGS	QLGEDYIQLH	
mouse	SSNAQLL LDY	CSSKGYNISW	ELGNEPNSFW	KKAHILIDGL	QLGEDFVELH	
rat	~~~~~	~~~~~	~~~~~	~~~~~	~~~~~	
						300
human	KLLRKSTFKN	AKLYGPDVGQ	PRRKTAKMLK	SFLKAGGEVI	DSVTWHHYYL	
mouse	KLLQRSAFQN	AKLYGPDIGQ	PRGKTVKLLR	SFLKAGGEVI	DSLTHHYYL	
rat	~~~~~	~~~~~	~~~~~	~~~~~	~~~~~	
						350
human	NGRTATREDF	LNPVDLIDIFI	SSVQKVFQVV	ESTRPGKKVW	LGETSSAYGG	
mouse	NGRIATKEDF	LSSDALDTFI	LSVQKILKVT	KEITPGKKVW	LGETSSAYGG	
rat	~~~~~	~~~~~	~~~~~	~~~~~	~~~~~	
						400
human	GAPLLSDTFA	AGFMWLDKLG	LSARMGIEVV	MRQVFFGAGN	YHLVDENFDP	
mouse	GAPLLSNTFA	AGFMWLDKLG	LSAQMGIIEVV	MRQVFFGAGN	YHLVDENFEP	
rat	~~~~~	~~~~~	~~~~~	~~~~~	~~~~~	
						450
human	LPDYWLSSLF	KKLVGTVLIM	ASVQGSKRRK	LRVYLNHCTNT	DNPRTYKEGDL	
mouse	LPDYWLSSLF	KKLVGPRVLL	SRVKGPDRSK	LRVYLNHCTNV	YHPRYQEGL	
rat	~~~~~	~~~~~	~~~~~	~~~~~	~~~~~	
						500
human	TLYAINLHNV	TKYLRLPYPF	SNKQVDKYLL	RPLGPHGLLS	KSVQLNGLTL	
mouse	TLYVLNLHNV	TKHLKVPPPL	FRKPVDTYLL	KPSGPDGLLS	KSVQLNGQIL	
rat	~~~~~	~~~~~	~~~~~	~~~~~	~~~~~	
						543
human	KMVDDQTLPP	LMEKPLRPGS	SLGLPAFFSYS	FFVIRNAKVA	ACI~	
mouse	KMVDEQTLPA	LTEKPLPAGS	ALSLPAFFSYG	FFVIRNAKIA	ACI~	
rat	KMVDEQTXPA	LTEKPLPAGS	SLSVPAFFSYG	FFVIRNAKIA	ACI~	

Fig. 17

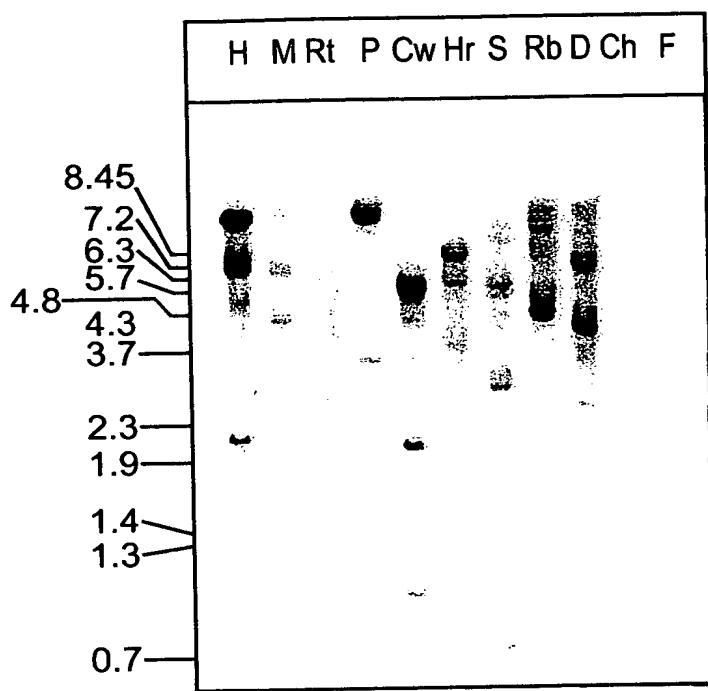


Fig. 18

Fig. 19